

SEQUENCE LISTING

<110> Soren Weis DAHL et al.
 <120> TRANSLOCATION DEPENDENT COMPLEMENTATION FOR DRUG SCREENING
 <130> 4614-0159PUS1
 <140> US 10/511,468
 <141> 2004-10-18
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 <170> PatentIn version 3.1
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 Gly Glu Gly Asp Ala Thr Tyr Gly Lys Leu Thr Leu Lys Phe Ile Cys
 35 40 45
 Thr Thr Gly Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr Phe
 50 55 60
 Ser Tyr Gly Val Gln Cys Phe Ser Arg Tyr Pro Asp His Met Lys Gln
 65 70 75 80
 His Asp Phe Phe Lys Ser Ala Met Pro Glu Gly Tyr Val Gln Glu Arg
 85 90 95
 Thr Ile Phe Phe Lys Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu Val
 100 105 110
 Lys Phe Glu Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly Ile
 115 120 125
 Asp Phe Lys Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn
 130 135 140
 Tyr Asn Ser His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly
 145 150 155 160
 Ile Lys Val Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Ser Val
 165 170 175
 Gln Leu Ala Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro

| 180 | | | | | 185 | | | | | 190 | | | | | |
|-------------------------|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Val | Leu | Leu | Pro | Asp | Asn | His | Tyr | Leu | Ser | Thr | Gln | Ser | Ala | Leu | Ser |
| | 195 | | | | | | 200 | | | | | 205 | | | |
| Lys | Asp | Pro | Asn | Glu | Lys | Arg | Asp | His | Met | Val | Leu | Leu | Glu | Phe | Val |
| | 210 | | | | | 215 | | | | | 220 | | | | |
| Thr | Ala | Ala | Gly | Ile | Thr | His | Gly | Met | Asp | Glu | Leu | Tyr | Lys | | |
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| <213> Aequorea victoria | | | | | | | | | | | | | | | |
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| Glu | Leu | Asp | Gly | Asp | Val | Asn | Gly | His | Lys | Phe | Ser | Val | Ser | Gly | Glu |
| | | | 20 | | | | | 25 | | | | | 30 | | |
| Gly | Glu | Gly | Asp | Ala | Thr | Tyr | Gly | Lys | Leu | Thr | Leu | Lys | Phe | Ile | Cys |
| | | 35 | | | | | 40 | | | | | 45 | | | |
| Thr | Thr | Gly | Lys | Leu | Pro | Val | Pro | Trp | Pro | Thr | Leu | Val | Thr | Thr | Phe |
| | 50 | | | | | 55 | | | | | 60 | | | | |
| Ser | Trp | Gly | Val | Gln | Cys | Phe | Ser | Arg | Tyr | Pro | Asp | His | Met | Lys | Gln |
| 65 | | | | 70 | | | | | | 75 | | | | | 80 |
| His | Asp | Phe | Phe | Lys | Ser | Ala | Met | Pro | Glu | Gly | Tyr | Val | Gln | Glu | Arg |
| | | | 85 | | | | | 90 | | | | | 95 | | |
| Thr | Ile | Phe | Phe | Lys | Asp | Asp | Gly | Asn | Tyr | Lys | Thr | Arg | Ala | Glu | Val |
| | | | 100 | | | | | 105 | | | | | 110 | | |
| Lys | Phe | Glu | Gly | Asp | Thr | Leu | Val | Asn | Arg | Ile | Glu | Leu | Lys | Gly | Ile |
| | | 115 | | | | | 120 | | | | | 125 | | | |
| Asp | Phe | Lys | Glu | Asp | Gly | Asn | Ile | Leu | Gly | His | Lys | Leu | Glu | Tyr | Asn |
| | 130 | | | | | 135 | | | | | 140 | | | | |
| Tyr | Asn | Ser | His | Asn | Val | Tyr | Ile | Met | Ala | Asp | Lys | Gln | Lys | Asn | Gly |
| 145 | | | | 150 | | | | | | 155 | | | | | 160 |
| Ile | Lys | Val | Asn | Phe | Lys | Ile | Arg | His | Asn | Ile | Glu | Asp | Gly | Ser | Val |
| | | | 165 | | | | | 170 | | | | | 175 | | |
| Gln | Leu | Ala | Asp | His | Tyr | Gln | Gln | Asn | Thr | Pro | Ile | Gly | Asp | Gly | Pro |
| | | 180 | | | | | | 185 | | | | 190 | | | |
| Val | Leu | Leu | Pro | Asp | Asn | His | Tyr | Leu | Ser | Thr | Gln | Ser | Ala | Leu | Ser |
| | 195 | | | | | | 200 | | | | | 205 | | | |
| Lys | Asp | Pro | Asn | Glu | Lys | Arg | Asp | His | Met | Val | Leu | Leu | Glu | Phe | Val |

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| | 20 | 25 30 |
| Gly Glu Gly Asp Ala Thr Tyr Gly Lys Leu Thr Leu Lys Phe Ile Cys | | |
| | 35 | 40 45 |
| Thr Thr Gly Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr Phe | | |
| | 50 | 55 60 |
| Ser His Gly Val Gln Cys Phe Ser Arg Tyr Pro Asp His Met Lys Gln | | |
| 65 | 70 | 75 80 |
| His Asp Phe Phe Lys Ser Ala Met Pro Glu Gly Tyr Val Gln Glu Arg | | |
| | 85 | 90 95 |
| Thr Ile Phe Phe Lys Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu Val | | |
| | 100 | 105 110 |
| Lys Phe Glu Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly Ile | | |
| | 115 | 120 125 |
| Asp Phe Lys Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn | | |
| | 130 | 135 140 |
| Tyr Asn Ser His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly | | |
| 145 | 150 | 155 160 |
| Ile Lys Val Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Ser Val | | |
| | 165 | 170 175 |
| Gln Leu Ala Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro | | |
| | 180 | 185 190 |
| Val Leu Leu Pro Asp Asn His Tyr Leu Ser Thr Gln Ser Ala Leu Ser | | |
| | 195 | 200 205 |
| Lys Asp Pro Asn Glu Lys Arg Asp His Met Val Leu Leu Glu Phe Val | | |
| | 210 | 215 220 |
| Thr Ala Ala Gly Ile Thr His Gly Met Asp Glu Leu Tyr Lys | | |
| 225 | 230 | 235 |

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<211> 239
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<400> 4

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Met Val Ser Lys Gly Glu Glu Leu Phe Thr Gly Val Val Pro Ile Leu
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Val Glu Leu Asp Gly Asp Val Asn Gly His Lys Phe Ser Val Ser Gly
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Glu Gly Glu Gly Asp Ala Thr Tyr Gly Lys Leu Thr Leu Lys Phe Ile
          35          40          45

Cys Thr Thr Gly Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr
          50          55          60

Leu Thr Tyr Gly Val Gln Cys Phe Ser Arg Tyr Pro Asp His Met Lys
65          70          75          80

Gln His Asp Phe Phe Lys Ser Ala Met Pro Glu Gly Tyr Val Gln Glu
          85          90          95

Arg Thr Ile Phe Phe Lys Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu
          100          105          110

Val Lys Phe Glu Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly
          115          120          125

Ile Asp Phe Lys Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr
          130          135          140

Asn Tyr Asn Ser His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn
145          150          155          160

Gly Ile Lys Val Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Ser
          165          170          175

Val Gln Leu Ala Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly
          180          185          190

Pro Val Leu Leu Pro Asp Asn His Tyr Leu Ser Thr Gln Ser Ala Leu
          195          200          205

Ser Lys Asp Pro Asn Glu Lys Arg Asp His Met Val Leu Leu Glu Phe
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Val Thr Ala Ala Gly Ile Thr Leu Gly Met Asp Glu Leu Tyr Lys
225          230          235

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<210> 5
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Met Val Ser Lys Gly Glu Glu Leu Phe Thr Gly Val Val Pro Ile Leu

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| Glu Gly Glu Gly Asp Ala Thr Tyr Gly Lys Leu Thr Leu Lys Phe Ile | 35 | 40 | 45 |
| Cys Thr Thr Gly Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr | 50 | 55 | 60 |
| Phe Gly Tyr Gly Leu Gln Cys Phe Ala Arg Tyr Pro Asp His Met Lys | 65 | 70 | 75 |
| Gln His Asp Phe Phe Lys Ser Ala Met Pro Glu Gly Tyr Val Gln Glu | 85 | 90 | 95 |
| Arg Thr Ile Phe Phe Lys Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu | 100 | 105 | 110 |
| Val Lys Phe Glu Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly | 115 | 120 | 125 |
| Ile Asp Phe Lys Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr | 130 | 135 | 140 |
| Asn Tyr Asn Ser His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn | 145 | 150 | 155 |
| Gly Ile Lys Val Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Ser | 165 | 170 | 175 |
| Val Gln Leu Ala Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly | 180 | 185 | 190 |
| Pro Val Leu Leu Pro Asp Asn His Tyr Leu Ser Tyr Gln Ser Ala Leu | 195 | 200 | 205 |
| Ser Lys Asp Pro Asn Glu Lys Arg Asp His Met Val Leu Leu Glu Phe | 210 | 215 | 220 |
| Val Thr Ala Ala Gly Ile Thr Leu Gly Met Asp Glu Leu Tyr Lys | 225 | 230 | 235 |

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|---|
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| Val Glu Leu Asp Gly Asp Val Asn Gly His Lys Phe Ser Val Ser Gly |
| 20 25 30 |
| Glu Gly Glu Gly Asp Ala Thr Tyr Gly Lys Leu Thr Leu Lys Phe Ile |

| 35 | 40 | 45 |
|--|----|----|
| Cys Thr Thr Gly Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr 50 55 60 | | |
| Leu Gly Tyr Gly Leu Gln Cys Phe Ala Arg Tyr Pro Asp His Met Lys 65 70 75 80 | | |
| Gln His Asp Phe Phe Lys Ser Ala Met Pro Glu Gly Tyr Val Gln Glu 85 90 95 | | |
| Arg Thr Ile Phe Phe Lys Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu 100 105 110 | | |
| Val Lys Phe Glu Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly 115 120 125 | | |
| Ile Asp Phe Lys Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr 130 135 140 | | |
| Asn Tyr Asn Ser His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn 145 150 155 160 | | |
| Gly Ile Lys Val Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Ser 165 170 175 | | |
| Val Gln Leu Ala Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly 180 185 190 | | |
| Pro Val Leu Leu Pro Asp Asn His Tyr Leu Ser Tyr Gln Ser Ala Leu 195 200 205 | | |
| Ser Lys Asp Pro Asn Glu Lys Arg Asp His Met Val Leu Leu Glu Phe 210 215 220 | | |
| Val Thr Ala Ala Gly Ile Thr Leu Gly Met Asp Glu Leu Tyr Lys 225 230 235 | | |

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| gcc aac aag aag gag ctg gcc cag ctg aag tgg gag ctg cag gcc ctg Ala Asn Lys Lys Glu Leu Ala Gln Leu Lys Trp Glu Leu Gln Ala Leu 20 25 30 | 95 |

aag aag gag ctg gcc cag tag gatcc
 Lys Lys Glu Leu Ala Gln
 35

121

<210> 8
 <211> 37
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 1 5 10 15
 Asn Lys Lys Glu Leu Ala Gln Leu Lys Trp Glu Leu Gln Ala Leu Lys
 20 25 30
 Lys Glu Leu Ala Gln
 35

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 1 5 10 15

47

aag ctg gcc cag ctg gag tgg aag aac cag gcc ctg gag aag aag ctg
 Lys Leu Ala Gln Leu Glu Trp Lys Asn Gln Ala Leu Glu Lys Lys Leu
 20 25 30

95

gcc cag ggc ggc acc ggt tag gatcc
 Ala Gln Gly Gly Thr Gly
 35

121

<210> 10
 <211> 37
 <212> PRT
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 1 5 10 15
 Leu Ala Gln Leu Glu Trp Lys Asn Gln Ala Leu Glu Lys Lys Leu Ala
 20 25 30
 Gln Gly Gly Thr Gly
 35

<210> 11
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1 5 10 15

Gly Ser Thr

<210> 12
<211> 19
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<213> Homo sapiens

<400> 12
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1 5 10 15

Gly Ser Thr

<210> 13
<211> 18
<212> PRT
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<400> 13
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1 5 10 15

Ser Gly

<210> 14
<211> 18
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<400> 14
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1 5 10 15

Ser Gly

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| <400> 16 | |
| gtcattccag ttttagaagc tc | 22 |
| <210> 17 | |
| <211> 28 | |
| <212> DNA | |
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| <220> | |
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| <210> 18 | |
| <211> 47 | |
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| <220> | |
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| <210> 19 | |
| <211> 38 | |
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| <213> Artificial Sequence | |
| <220> | |
| <223> Synthetic oligonucleotide primer sequence | |
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| <211> 41 | |
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 catg 64

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 <212> DNA
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 <210> 44
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gggc 60
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<400> 48
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<210> 49

<211> 29
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<210> 58
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 <213> Artificial Sequence

<220>
 <223> Synthetic oligonucleotide primer sequence
 <400> 61
 ccaccatgcc gcagctgaac ggcgggtgga 29

<210> 62
 <211> 22
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Synthetic oligonucleotide primer sequence
 <400> 62
 caggcggagg ccgtctttcc gc 22

<210> 63
 <211> 36

<212> DNA
 <213> Artificial Sequence

 <220>
 <223> Synthetic oligonucleotide primer sequence

 <400> 63
 gttcagctga tgggagtgca ggtggaaacc atctcc 36

<210> 64
 <211> 60
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Synthetic oligonucleotide primer sequence

 <400> 64
 gttggatccg atatcaccgg tgccgtgtccc agttccttcc agttttagaa gctccacatc 60

<210> 65
 <211> 62
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Synthetic oligonucleotide primer sequence

 <400> 65
 gttgctagcg atatccggaa cgggcactgg gaccgggtgag atgtggcatg aaggcctgga 60
 ag 62

<210> 66
 <211> 35
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Synthetic oligonucleotide primer sequence

 <400> 66
 gttcccggga gctgctttga gattcgtcgg aacac 35

<210> 67
 <211> 29
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Synthetic oligonucleotide primer sequence

 <400> 67
 ccaccatgcc tgagccagcc aagtctgct 29

<210> 68
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic oligonucleotide primer sequence

<400> 68
ccttgaggagct ggtgtacttg gtagac

25

<210> 69
<211> 18
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic peptide linker sequence

<400> 69
Asp Pro Ala Phe Leu Tyr Lys Val Val Ile Ser Gly Ser Gly Ser Gly
1 5 10 15

Ser Gly

<210> 70
<211> 19
<212> PRT
<213> Artificial sequence

<220>
<223> Synthetic peptide linker sequence

<400> 70
Gly Ser Gly Ser Gly Ser Gly Asp Ile Thr Ser Leu Tyr Lys Lys Ala
1 5 10 15

Gly Ser Thr

<210> 71
<211> 16
<212> PRT
<213> Artificial sequence

<220>
<223> Synthetic peptide linker sequence

<400> 71
Asp Pro Ala Phe Leu Tyr Lys Val Val Ile Ser Gly Ser Gly Ser Gly
1 5 10 15

<210> 72
<211> 9

<212> PRT
 <213> Artificial sequence

 <220>
 <223> Synthetic peptide linker sequence

 <400> 72
 Gly Ser Gly Ser Gly Ser Gly Asp Leu
 1 5

 <210> 73
 <211> 18
 <212> PRT
 <213> Artificial sequence

 <220>
 <223> Synthetic peptide linker sequence

 <400> 73
 Asp Pro Ala Phe Leu Tyr Lys Val Val Ile Ser Gly Thr Gly Thr Gly
 1 5 10 15

 Thr Gly

 <210> 74
 <211> 19
 <212> PRT
 <213> Artificial sequence

 <220>
 <223> Synthetic peptide linker sequence

 <400> 74
 Gly Ser Gly Ser Gly Ser Gly Asp Ile Thr Ser Leu Tyr Lys Lys Ala
 1 5 10 15

 Gly Ser Thr

 <210> 75
 <211> 12
 <212> PRT
 <213> Artificial sequence

 <220>
 <223> Synthetic peptide linker sequence

 <400> 75
 Asp Ile Thr Ser Leu Tyr Lys Lys Ala Gly Ser Thr
 1 5 10

 <210> 76
 <211> 19
 <212> PRT
 <213> Artificial sequence

<220>
 <223> Synthetic peptide linker sequence

 <400> 76
 Gly Thr Gly Thr Gly Thr Gly Asp Ile Thr Ser Leu Tyr Lys Lys Ala
 1 5 10 15

 Gly Ser Thr

 <210> 77
 <211> 10
 <212> PRT
 <213> Artificial sequence

 <220>
 <223> Synthetic peptide linker sequence

 <400> 77
 Leu Pro Ser Gly Ser Gly Ser Gly Ser Gly
 1 5 10

 <210> 78
 <211> 9
 <212> PRT
 <213> Aequorea victoria

 <400> 78

 Leu Thr Tyr Gly Val Gln Cys Phe Ser
 1 5

 <210> 79
 <211> 9
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> synthetically mutated peptide L64F:T65G:V68L:S72A

 <400> 79

 Phe Gly Tyr Gly Leu Gln Cys Phe Ala
 1 5

 <210> 80
 <211> 9
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> synthetically mutated peptide T65G:V68L:S72A

 <400> 80

Leu Gly Tyr Gly Leu Gln Cys Phe Ala
1 5

<210> 81
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic peptide linker sequence

<400> 81

Gly Ser Gly Ser Gly Ser Gly
1 5

<210> 82
<211> 6
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic peptide linker sequence

<400> 82

Gly Gly Thr Gly Ser Gly
1 5

<210> 83
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetic peptide linker sequence

<400> 83

Gly Thr Gly Thr Gly Thr Gly
1 5

<210> 84
<211> 238
<212> PRT
<213> Aequorea Victoria

<400> 84

Met Ser Lys Gly Glu Glu Leu Phe Thr Gly Val Val Pro Ile Leu Val
1 5 10 15

Glu Leu Asp Gly Asp Val Asn Gly His Lys Phe Ser Val Ser Gly Glu
20 25 30

Gly Glu Gly Asp Ala Thr Tyr Gly Lys Leu Thr Leu Lys Phe Ile Cys
35 40 45

Thr Thr Gly Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr Phe
50 55 60

Ser Tyr Gly Val Gln Cys Phe Ser Arg Tyr Pro Asp His Met Lys Gln
65 70 75 80

His Asp Phe Phe Lys Ser Ala Met Pro Glu Gly Tyr Val Gln Glu Arg
85 90 95

Thr Ile Phe Phe Lys Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu Val
100 105 110

Lys Phe Glu Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly Ile
115 120 125

Asp Phe Lys Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn
130 135 140

Tyr Asn Ser His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly
145 150 155 160

Ile Lys Val Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Ser Val
165 170 175

Gln Leu Ala Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro
180 185 190

Val Leu Leu Pro Asp Asn His Tyr Leu Ser Thr Gln Ser Ala Leu Ser
195 200 205

Lys Asp Pro Asn Glu Lys Arg Asp His Met Val Leu Leu Glu Phe Val
210 215 220

Thr Ala Ala Gly Ile Thr His Gly Met Asp Glu Leu Tyr Lys
225 230 235

<210> 85

<211> 238
 <212> PRT
 <213> Renilla muelleri

<400> 85

Met Ser Lys Gln Ile Leu Lys Asn Thr Cys Leu Gln Glu Val Met Ser
 1 5 10 15

Tyr Lys Val Asn Leu Glu Gly Ile Val Asn Asn His Val Phe Thr Met
 20 25 30

Glu Gly Cys Gly Lys Gly Asn Ile Leu Phe Gly Asn Gln Leu Val Gln
 35 40 45

Ile Arg Val Thr Lys Gly Ala Pro Leu Pro Phe Ala Phe Asp Ile Val
 50 55 60

Ser Pro Ala Phe Gln Tyr Gly Asn Arg Thr Phe Thr Lys Tyr Pro Asn
 65 70 75 80

Asp Ile Ser Asp Tyr Phe Ile Gln Ser Phe Pro Ala Gly Phe Met Tyr
 85 90 95

Glu Arg Thr Leu Arg Tyr Glu Asp Gly Gly Leu Val Glu Ile Arg Ser
 100 105 110

Asp Ile Asn Leu Ile Glu Asp Lys Phe Val Tyr Arg Val Glu Tyr Lys
 115 120 125

Gly Ser Asn Phe Pro Asp Asp Gly Pro Val Met Gln Lys Thr Ile Leu
 130 135 140

Gly Ile Glu Pro Ser Phe Glu Ala Met Tyr Met Asn Asn Gly Val Leu
 145 150 155 160

Val Gly Glu Val Ile Leu Val Tyr Lys Leu Asn Ser Gly Lys Tyr Tyr
 165 170 175

Ser Cys His Met Lys Thr Leu Met Lys Ser Lys Gly Val Val Lys Glu
 180 185 190

Phe Pro Ser Tyr His Phe Ile Gln His Arg Leu Glu Lys Thr Tyr Val
 195 200 205

Glu Asp Gly Gly Phe Val Glu Gln His Glu Thr Ala Ile Ala Gln Met
 210 215 220

Thr Ser Ile Gly Lys Pro Leu Gly Ser Leu His Glu Trp Val
 225 230 235

<210> 86
 <211> 233
 <212> PRT
 <213> Renilla reniformis

<400> 86

Met Asp Leu Ala Lys Leu Gly Leu Lys Glu Val Met Pro Thr Lys Ile
 1 5 10 15

Asn Leu Glu Gly Leu Val Gly Asp His Ala Phe Ser Met Glu Gly Val
 20 25 30

Gly Glu Gly Asn Ile Leu Glu Gly Thr Gln Glu Val Lys Ile Ser Val
 35 40 45

Thr Lys Gly Ala Pro Leu Pro Phe Ala Phe Asp Ile Val Ser Val Ala
 50 55 60

Phe Ser Tyr Gly Asn Arg Ala Tyr Thr Gly Tyr Pro Glu Glu Ile Ser
 65 70 75 80

Asp Tyr Phe Leu Gln Ser Phe Pro Glu Gly Phe Thr Tyr Glu Arg Asn
 85 90 95

Ile Arg Tyr Gln Asp Gly Gly Thr Ala Ile Val Lys Ser Asp Ile Ser
 100 105 110

Leu Glu Asp Gly Lys Phe Ile Val Asn Val Asp Phe Lys Ala Lys Asp
 115 120 125

Leu Arg Arg Met Gly Pro Val Met Gln Gln Asp Ile Val Gly Met Gln
 130 135 140

Pro Ser Tyr Glu Ser Met Tyr Thr Asn Val Thr Ser Val Ile Gly Glu
 145 150 155 160

Cys Ile Ile Ala Phe Lys Leu Gln Thr Gly Lys His Phe Thr Tyr His

| | | | | | |
|---|-----|---|-----|--|-----|
| | 165 | | 170 | | 175 |
| Met Arg Thr Val Tyr Lys Ser Lys Lys Pro Val Glu Thr Met Pro Leu | 180 | | 185 | | 190 |
| Tyr His Phe Ile Gln His Arg Leu Val Lys Thr Asn Val Asp Thr Ala | 195 | | 200 | | 205 |
| Ser Gly Tyr Val Val Gln His Glu Thr Ala Ile Ala Ala His Ser Thr | 210 | | 215 | | 220 |
| Ile Lys Lys Ile Glu Gly Ser Leu Pro | 225 | | 230 | | |
| <210> 87 | | | | | |
| <211> 225 | | | | | |
| <212> PRT | | | | | |
| <213> Discosoma sp. | | | | | |
| <400> 87 | | | | | |
| Met Arg Ser Ser Lys Asn Val Ile Lys Glu Phe Met Arg Phe Lys Val | 1 | 5 | 10 | | 15 |
| Arg Met Glu Gly Thr Val Asn Gly His Glu Phe Glu Ile Glu Gly Glu | 20 | | 25 | | 30 |
| Gly Glu Gly Arg Pro Tyr Glu Gly His Asn Thr Val Lys Leu Lys Val | 35 | | 40 | | 45 |
| Thr Lys Gly Gly Pro Leu Pro Phe Ala Trp Asp Ile Leu Ser Pro Gln | 50 | | 55 | | 60 |
| Phe Gln Tyr Gly Ser Lys Val Tyr Val Lys His Pro Ala Asp Ile Pro | 65 | | 70 | | 75 |
| | | | | | 80 |
| Asp Tyr Lys Lys Leu Ser Phe Pro Glu Gly Phe Lys Trp Glu Arg Val | 85 | | 90 | | 95 |
| Met Asn Phe Glu Asp Gly Gly Val Val Thr Val Thr Gln Asp Ser Ser | 100 | | 105 | | 110 |
| Leu Gln Asp Gly Cys Phe Ile Tyr Lys Val Lys Phe Ile Gly Val Asn | 115 | | 120 | | 125 |

Phe Pro Ser Asp Gly Pro Val Met Gln Lys Lys Thr Met Gly Trp Glu
 130 135 140

Ala Ser Thr Glu Arg Leu Tyr Pro Arg Asp Gly Val Leu Lys Gly Glu
 145 150 155 160

Ile His Lys Ala Leu Lys Leu Lys Asp Gly Gly His Tyr Leu Val Glu
 165 170 175

Phe Lys Ser Ile Tyr Met Ala Lys Lys Pro Val Gln Leu Pro Gly Tyr
 180 185 190

Tyr Tyr Val Asp Ser Lys Leu Asp Ile Thr Ser His Asn Glu Asp Tyr
 195 200 205

Thr Ile Val Glu Gln Tyr Glu Arg Thr Glu Gly Arg His His Leu Phe
 210 215 220

Leu
 225

<210> 88
 <211> 148
 <212> PRT
 <213> Anemonia sulcata

<400> 88

Met Ala Ser Phe Leu Lys Lys Thr Met Pro Phe Lys Thr Thr Ile Glu
 1 5 10 15

Gly Thr Val Asn Gly His Tyr Phe Lys Cys Thr Gly Lys Gly Glu Gly
 20 25 30

Asn Pro Phe Glu Gly Thr Gln Glu Met Lys Ile Glu Val Ile Glu Gly
 35 40 45

Gly Pro Leu Pro Phe Ala Phe His Ile Leu Ser Thr Ser Cys Met Tyr
 50 55 60

Gly Ser Lys Thr Phe Ile Lys Tyr Val Ser Gly Ile Pro Asp Tyr Phe
 65 70 75 80

Lys Gln Ser Phe Pro Glu Gly Phe Thr Trp Glu Arg Thr Thr Thr Tyr

85

90

95

Glu Asp Gly Gly Phe Leu Thr Ala His Gln Asp Thr Ser Leu Asp Gly
 100 105 110

Asp Cys Leu Val Tyr Lys Val Lys Ile Leu Gly Asn Asn Phe Pro Ala
 115 120 125

Asp Gly Pro Arg Asp Ala Glu Gln Ser Arg Lys Met Gly Ala Ser His
 130 135 140

Arg Asp Thr Leu
 145

<210> 89
 <211> 231
 <212> PRT
 <213> Zoanthus sp.

<400> 89

Met Ala Gln Ser Lys His Gly Leu Thr Lys Glu Met Thr Met Lys Tyr
 1 5 10 15

Arg Met Glu Gly Cys Val Asp Gly His Lys Phe Val Ile Thr Gly Glu
 20 25 30

Gly Ile Gly Tyr Pro Phe Lys Gly Lys Gln Ala Ile Asn Leu Cys Val
 35 40 45

Val Glu Gly Gly Pro Leu Pro Phe Ala Glu Asp Ile Leu Ser Ala Ala
 50 55 60

Phe Asn Tyr Gly Asn Arg Val Phe Thr Glu Tyr Pro Gln Asp Ile Val
 65 70 75 80

Asp Tyr Phe Lys Asn Ser Cys Pro Ala Gly Tyr Thr Trp Asp Arg Ser
 85 90 95

Phe Leu Phe Glu Asp Gly Ala Val Cys Ile Cys Asn Ala Asp Ile Thr
 100 105 110

Val Ser Val Glu Glu Asn Cys Met Tyr His Glu Ser Lys Phe Tyr Gly
 115 120 125

Val Asn Phe Pro Ala Asp Gly Pro Val Met Lys Lys Met Thr Asp Asn
 130 135 140

Trp Glu Pro Ser Cys Glu Lys Ile Ile Pro Val Pro Lys Gln Gly Ile
 145 150 155 160

Leu Lys Gly Asp Val Ser Met Tyr Leu Leu Leu Lys Asp Gly Gly Arg
 165 170 175

Leu Arg Cys Gln Phe Asp Thr Val Tyr Lys Ala Lys Ser Val Pro Arg
 180 185 190

Lys Met Pro Asp Trp His Phe Ile Gln His Lys Leu Thr Arg Glu Asp
 195 200 205

Arg Ser Asp Ala Lys Asn Gln Lys Trp His Leu Thr Glu His Ala Ile
 210 215 220

Ala Ser Gly Ser Ala Leu Pro
 225 230

<210> 90
 <211> 232
 <212> PRT
 <213> Anemonia sulcata

<400> 90

Met Ala Ser Phe Leu Lys Lys Thr Met Pro Phe Lys Thr Thr Ile Glu
 1 5 10 15

Gly Thr Val Asn Gly His Tyr Phe Lys Cys Thr Gly Lys Gly Glu Gly
 20 25 30

Asn Pro Phe Glu Gly Thr Gln Glu Met Lys Ile Glu Val Ile Glu Gly
 35 40 45

Gly Pro Leu Pro Phe Ala Phe His Ile Leu Ser Thr Ser Cys Met Tyr
 50 55 60

Gly Ser Lys Thr Phe Ile Lys Tyr Val Ser Gly Ile Pro Asp Tyr Phe
 65 70 75 80

Lys Gln Ser Phe Pro Glu Gly Phe Thr Trp Glu Arg Thr Thr Thr Tyr

| | | | | | |
|---|-----|--|-----|--|-----|
| | 85 | | 90 | | 95 |
| Glu Asp Gly Gly Phe Leu Thr Ala His Gln Asp Thr Ser Leu Asp Gly | 100 | | 105 | | 110 |
| Asp Cys Leu Val Tyr Lys Val Lys Ile Leu Gly Asn Asn Phe Pro Ala | 115 | | 120 | | 125 |
| Asp Gly Pro Val Met Gln Asn Lys Ala Gly Arg Trp Glu Pro Ala Thr | 130 | | 135 | | 140 |
| Glu Ile Val Tyr Glu Val Asp Gly Val Leu Arg Gly Gln Ser Leu Met | 145 | | 150 | | 155 |
| | | | | | 160 |
| Ala Leu Lys Cys Pro Gly Gly Arg His Leu Thr Cys His Leu His Thr | 165 | | 170 | | 175 |
| Thr Tyr Arg Ser Lys Lys Pro Ala Ser Ala Leu Lys Met Pro Gly Phe | 180 | | 185 | | 190 |
| His Phe Glu Asp His Arg Ile Glu Ile Met Glu Glu Val Glu Lys Gly | 195 | | 200 | | 205 |
| Lys Cys Tyr Lys Gln Tyr Glu Ala Ala Val Gly Arg Tyr Cys Asp Ala | 210 | | 215 | | 220 |
| Ala Pro Ser Lys Leu Gly His Asn | 225 | | 230 | | |

<210> 91
 <211> 228
 <212> PRT
 <213> Anemonia sulcata

<400> 91

| | | | |
|---|---|----|----|
| Met Tyr Pro Ser Ile Lys Glu Thr Met Arg Val Gln Leu Ser Met Glu | | | |
| 1 | 5 | 10 | 15 |

| | | | |
|---|----|----|--|
| Gly Ser Val Asn Tyr His Ala Phe Lys Cys Thr Gly Lys Gly Glu Gly | | | |
| 20 | 25 | 30 | |

| | | | |
|---|----|----|--|
| Lys Pro Tyr Glu Gly Thr Gln Ser Leu Asn Ile Thr Ile Thr Glu Gly | | | |
| 35 | 40 | 45 | |

Gly Pro Leu Pro Phe Ala Phe Asp Ile Leu Ser His Ala Phe Gln Tyr
50 55 60

Gly Ile Lys Val Phe Ala Lys Tyr Pro Lys Glu Ile Pro Asp Phe Phe
65 70 75 80

Lys Gln Ser Leu Pro Gly Gly Phe Ser Trp Glu Arg Val Ser Thr Tyr
85 90 95

Glu Asp Gly Gly Val Leu Ser Ala Thr Gln Glu Thr Ser Leu Gln Gly
100 105 110

Asp Cys Ile Ile Cys Lys Val Lys Val Leu Gly Thr Asn Phe Pro Ala
115 120 125

Asn Gly Pro Val Met Gln Lys Lys Thr Cys Gly Trp Glu Pro Ser Thr
130 135 140

Glu Thr Val Ile Pro Arg Asp Gly Gly Leu Leu Leu Arg Asp Thr Pro
145 150 155 160

Ala Leu Met Leu Ala Asp Gly Gly His Leu Ser Cys Phe Met Glu Thr
165 170 175

Thr Tyr Lys Ser Lys Lys Glu Val Lys Leu Pro Glu Leu His Phe His
180 185 190

His Leu Arg Met Glu Lys Leu Asn Ile Ser Asp Asp Trp Lys Thr Val
195 200 205

Glu Gln His Glu Ser Val Val Ala Ser Tyr Ser Gln Val Pro Ser Lys
210 215 220

Leu Gly His Asn
225

<210> 92
<211> 225
<212> PRT
<213> Montastraea cavernosa

<400> 92

Met Ser Val Ile Lys Pro Ile Met Glu Ile Lys Leu Arg Met Gln Gly

| | | | |
|---|-----|-----|-----|
| 1 | 5 | 10 | 15 |
| Val Val Asn Gly His Lys Phe Val Ile Lys Gly Glu Gly Glu Gly Lys | 20 | 25 | 30 |
| Pro Phe Glu Gly Thr Gln Thr Ile Asn Leu Thr Val Lys Glu Gly Ala | 35 | 40 | 45 |
| Pro Leu Pro Phe Ala Tyr Asp Ile Leu Thr Ser Ala Phe Gln Tyr Gly | 50 | 55 | 60 |
| Asn Arg Val Phe Thr Lys Tyr Pro Asp Asp Ile Pro Asp Tyr Phe Lys | 65 | 70 | 75 |
| Gln Thr Phe Pro Glu Gly Tyr Ser Trp Glu Arg Ile Met Ala Tyr Glu | 85 | 90 | 95 |
| Asp Gln Ser Ile Cys Thr Ala Thr Ser Asp Ile Lys Met Glu Gly Asp | 100 | 105 | 110 |
| Cys Phe Ile Tyr Glu Ile Gln Phe His Gly Val Asn Phe Pro Pro Asn | 115 | 120 | 125 |
| Gly Pro Val Met Gln Lys Lys Thr Leu Lys Trp Glu Pro Ser Thr Glu | 130 | 135 | 140 |
| Lys Met Tyr Val Arg Asp Gly Val Leu Lys Gly Asp Val Asn Met Ala | 145 | 150 | 155 |
| Leu Leu Leu Glu Gly Gly Gly His Tyr Arg Cys Asp Phe Arg Ser Thr | 165 | 170 | 175 |
| Tyr Lys Ala Lys Lys Arg Val Gln Leu Pro Asp Tyr His Phe Val Asp | 180 | 185 | 190 |
| His Arg Ile Glu Ile Leu Ser His Asp Asn Asp Tyr Asn Thr Val Lys | 195 | 200 | 205 |
| Leu Ser Glu Asp Ala Glu Ala Arg Tyr Ser Met Leu Pro Ser Gln Ala | 210 | 215 | 220 |
| Lys | | | |
| 225 | | | |

<210> 93
 <211> 227
 <212> PRT
 <213> Montastraea faveolata

<400> 93

Met Ser Val Ile Lys Pro Asp Met Lys Ile Lys Leu Arg Met Glu Gly
 1 5 10 15

Ala Val Asn Gly His Lys Phe Val Ile Glu Gly Asp Gly Lys Gly Lys
 20 25 30

Pro Phe Glu Gly Thr Gln Ser Met Asp Leu Thr Val Lys Glu Gly Ala
 35 40 45

Pro Leu Pro Phe Ala Tyr Asp Ile Leu Thr Thr Val Phe Asp Tyr Gly
 50 55 60

Asn Arg Val Phe Ala Lys Tyr Pro Gln Asp Ile Pro Asp Tyr Phe Lys
 65 70 75 80

Gln Thr Phe Pro Glu Gly Tyr Ser Trp Glu Arg Ser Met Thr Tyr Glu
 85 90 95

Asp Gln Gly Ile Cys Val Ala Thr Asn Asp Ile Thr Leu Met Lys Gly
 100 105 110

Val Asp Asp Cys Phe Val Tyr Lys Ile Arg Phe Asp Gly Val Asn Phe
 115 120 125

Pro Ala Asn Gly Pro Val Met Gln Lys Lys Thr Leu Lys Trp Glu Pro
 130 135 140

Ser Thr Glu Lys Met Tyr Val Arg Asp Gly Val Leu Lys Gly Asp Val
 145 150 155 160

Asn Met Ala Leu Leu Leu Glu Gly Gly Gly His Tyr Arg Cys Asp Phe
 165 170 175

Lys Thr Thr Tyr Lys Ala Lys Lys Phe Val Gln Leu Pro Asp Tyr His
 180 185 190

Phe Val Asp His Arg Ile Glu Ile Leu Ser His Asp Lys Asp Tyr Asn
 195 200 205

Lys Val Lys Leu Tyr Glu His Ala Glu Ala His Ser Gly Leu Pro Arg
 210 215 220

Gln Ala Lys
 225

<210> 94
 <211> 266
 <212> PRT
 <213> Clavularia sp.

<400> 94

Met Lys Cys Lys Phe Val Phe Cys Leu Ser Phe Leu Val Leu Ala Ile
 1 5 10 15

Thr Asn Ala Asn Ile Phe Leu Arg Asn Glu Ala Asp Leu Glu Glu Lys
 20 25 30

Thr Leu Arg Ile Pro Lys Ala Leu Thr Thr Met Gly Val Ile Lys Pro
 35 40 45

Asp Met Lys Ile Lys Leu Lys Met Glu Gly Asn Val Asn Gly His Ala
 50 55 60

Phe Val Ile Glu Gly Glu Gly Glu Gly Lys Pro Tyr Asp Gly Thr His
 65 70 75 80

Thr Leu Asn Leu Glu Val Lys Glu Gly Ala Pro Leu Pro Phe Ser Tyr
 85 90 95

Asp Ile Leu Ser Asn Ala Phe Gln Tyr Gly Asn Arg Ala Leu Thr Lys
 100 105 110

Tyr Pro Asp Asp Ile Ala Asp Tyr Phe Lys Gln Ser Phe Pro Glu Gly
 115 120 125

Tyr Ser Trp Glu Arg Thr Met Thr Phe Glu Asp Lys Gly Ile Val Lys
 130 135 140

Val Lys Ser Asp Ile Ser Met Glu Glu Asp Ser Phe Ile Tyr Glu Ile
 145 150 155 160

Arg Phe Asp Gly Met Asn Phe Pro Pro Asn Gly Pro Val Met Gln Lys
 165 170 175

Lys Thr Leu Lys Trp Glu Pro Ser Thr Glu Ile Met Tyr Val Arg Asp
 180 185 190

Gly Val Leu Val Gly Asp Ile Ser His Ser Leu Leu Leu Glu Gly Gly
 195 200 205

Gly His Tyr Arg Cys Asp Phe Lys Ser Ile Tyr Lys Ala Lys Lys Val
 210 215 220

Val Lys Leu Pro Asp Tyr His Phe Val Asp His Arg Ile Glu Ile Leu
 225 230 235 240

Asn His Asp Lys Asp Tyr Asn Lys Val Thr Leu Tyr Glu Asn Ala Val
 245 250 255

Ala Arg Tyr Ser Leu Leu Pro Ser Gln Ala
 260 265

<210> 95
 <211> 232
 <212> PRT
 <213> Discosoma striata

<400> 95

Met Ser Cys Ser Lys Ser Val Ile Lys Glu Glu Met Leu Ile Asp Leu
 1 5 10 15

His Leu Glu Gly Thr Phe Asn Gly His Tyr Phe Glu Ile Lys Gly Lys
 20 25 30

Gly Lys Gly Gln Pro Asn Glu Gly Thr Asn Thr Val Thr Leu Glu Val
 35 40 45

Thr Lys Gly Gly Pro Leu Pro Phe Gly Trp His Ile Leu Cys Pro Gln
 50 55 60

Phe Gln Tyr Gly Asn Lys Ala Phe Val His His Pro Asp Asn Ile His
 65 70 75 80

Asp Tyr Leu Lys Leu Ser Phe Pro Glu Gly Tyr Thr Trp Glu Arg Ser
85 90 95

Met His Phe Glu Asp Gly Gly Leu Cys Cys Ile Thr Asn Asp Ile Ser
100 105 110

Leu Thr Gly Asn Cys Phe Tyr Tyr Asp Ile Lys Phe Thr Gly Leu Asn
115 120 125

Phe Pro Pro Asn Gly Pro Val Val Gln Lys Lys Thr Thr Gly Trp Glu
130 135 140

Pro Ser Thr Glu Arg Leu Tyr Pro Arg Asp Gly Val Leu Ile Gly Asp
145 150 155 160

Ile His His Ala Leu Thr Val Glu Gly Gly Gly His Tyr Ala Cys Asp
165 170 175

Ile Lys Thr Val Tyr Arg Ala Lys Lys Ala Ala Leu Lys Met Pro Gly
180 185 190

Tyr His Tyr Val Asp Thr Lys Leu Val Ile Trp Asn Asn Asp Lys Glu
195 200 205

Phe Met Lys Val Glu Glu His Glu Ile Ala Val Ala Arg His His Pro
210 215 220

Phe Tyr Glu Pro Lys Lys Asp Lys
225 230

<210> 96
<211> 238
<212> PRT
<213> Ptilosarcus sp.

<400> 96

Met Asn Arg Asn Val Leu Lys Asn Thr Gly Leu Lys Glu Ile Met Ser
1 5 10 15

Ala Lys Ala Ser Val Glu Gly Ile Val Asn Asn His Val Phe Ser Met
20 25 30

Glu Gly Phe Gly Lys Gly Asn Val Leu Phe Gly Asn Gln Leu Met Gln
35 40 45

Ile Arg Val Thr Lys Gly Gly Pro Leu Pro Phe Ala Phe Asp Ile Val
50 55 60

Ser Ile Ala Phe Gln Tyr Gly Asn Arg Thr Phe Thr Lys Tyr Pro Asp
65 70 75 80

Asp Ile Ala Asp Tyr Phe Val Gln Ser Phe Pro Ala Gly Phe Phe Tyr
85 90 95

Glu Arg Asn Leu Arg Phe Glu Asp Gly Ala Ile Val Asp Ile Arg Ser
100 105 110

Asp Ile Ser Leu Glu Asp Asp Lys Phe His Tyr Lys Val Glu Tyr Arg
115 120 125

Gly Asn Gly Phe Pro Ser Asn Gly Pro Val Met Gln Lys Ala Ile Leu
130 135 140

Gly Met Glu Pro Ser Phe Glu Val Val Tyr Met Asn Ser Gly Val Leu
145 150 155 160

Val Gly Glu Val Asp Leu Val Tyr Lys Leu Glu Ser Gly Asn Tyr Tyr
165 170 175

Ser Cys His Met Lys Thr Phe Tyr Arg Ser Lys Gly Gly Val Lys Glu
180 185 190

Phe Pro Glu Tyr His Phe Ile His His Arg Leu Glu Lys Thr Tyr Val
195 200 205

Glu Glu Gly Ser Phe Val Glu Gln His Glu Thr Ala Ile Ala Gln Leu
210 215 220

Thr Thr Ile Gly Lys Pro Leu Gly Ser Leu His Glu Trp Val
225 230 235

<210> 97
<211> 231
<212> PRT
<213> Zoanthus sp.

<400> 97

Met Ala His Ser Lys His Gly Leu Lys Glu Glu Met Thr Met Lys Tyr
 1 5 10 15
 His Met Glu Gly Cys Val Asn Gly His Lys Phe Val Ile Thr Gly Glu
 20 25 30
 Gly Ile Gly Tyr Pro Phe Lys Gly Lys Gln Thr Ile Asn Leu Cys Val
 35 40 45
 Ile Glu Gly Gly Pro Leu Pro Phe Ser Glu Asp Ile Leu Ser Ala Gly
 50 55 60
 Phe Lys Tyr Gly Asp Arg Ile Phe Thr Glu Tyr Pro Gln Asp Ile Val
 65 70 75 80
 Asp Tyr Phe Lys Asn Ser Cys Pro Ala Gly Tyr Thr Trp Gly Arg Ser
 85 90 95
 Phe Leu Phe Glu Asp Gly Ala Val Cys Ile Cys Asn Val Asp Ile Thr
 100 105 110
 Val Ser Val Lys Glu Asn Cys Ile Tyr His Lys Ser Ile Phe Asn Gly
 115 120 125
 Met Asn Phe Pro Ala Asp Gly Pro Val Met Lys Lys Met Thr Thr Asn
 130 135 140
 Trp Glu Ala Ser Cys Glu Lys Ile Met Pro Val Pro Lys Gln Gly Ile
 145 150 155 160
 Leu Lys Gly Asp Val Ser Met Tyr Leu Leu Leu Lys Asp Gly Gly Arg
 165 170 175
 Tyr Arg Cys Gln Phe Asp Thr Val Tyr Lys Ala Lys Ser Val Pro Ser
 180 185 190
 Lys Met Pro Glu Trp His Phe Ile Gln His Lys Leu Leu Arg Glu Asp
 195 200 205
 Arg Ser Asp Ala Lys Asn Gln Lys Trp Gln Leu Thr Glu His Ala Ile
 210 215 220
 Ala Phe Pro Ser Ala Leu Ala

225

230

<210> 98
 <211> 229
 <212> PRT
 <213> Anemonia majano

<400> 98

Met Ala Leu Ser Asn Lys Phe Ile Gly Asp Asp Met Lys Met Thr Tyr
 1 5 10 15

His Met Asp Gly Cys Val Asn Gly His Tyr Phe Thr Val Lys Gly Glu
 20 25 30

Gly Asn Gly Lys Pro Tyr Glu Gly Thr Gln Thr Ser Thr Phe Lys Val
 35 40 45

Thr Met Ala Asn Gly Gly Pro Leu Ala Phe Ser Phe Asp Ile Leu Ser
 50 55 60

Thr Val Phe Lys Tyr Gly Asn Arg Cys Phe Thr Ala Tyr Pro Thr Ser
 65 70 75 80

Met Pro Asp Tyr Phe Lys Gln Ala Phe Pro Asp Gly Met Ser Tyr Glu
 85 90 95

Arg Thr Phe Thr Tyr Glu Asp Gly Gly Val Ala Thr Ala Ser Trp Glu
 100 105 110

Ile Ser Leu Lys Gly Asn Cys Phe Glu His Lys Ser Thr Phe His Gly
 115 120 125

Val Asn Phe Pro Ala Asp Gly Pro Val Met Ala Lys Lys Thr Thr Gly
 130 135 140

Trp Asp Pro Ser Phe Glu Lys Met Thr Val Cys Asp Gly Ile Leu Lys
 145 150 155 160

Gly Asp Val Thr Ala Phe Leu Met Leu Gln Gly Gly Gly Asn Tyr Arg
 165 170 175

Cys Gln Phe His Thr Ser Tyr Lys Thr Lys Lys Pro Val Thr Met Pro
 180 185 190

Pro Asn His Val Val Glu His Arg Ile Ala Arg Thr Asp Leu Asp Lys
 195 200 205

Gly Gly Asn Ser Val Gln Leu Thr Glu His Ala Val Ala His Ile Thr
 210 215 220

Ser Val Val Pro Phe
 225

<210> 99
 <211> 238
 <212> PRT
 <213> Aequorea macrodactyla

<400> 99

Met Ser Lys Gly Glu Glu Leu Phe Thr Gly Ile Val Pro Val Leu Ile
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Glu Leu Asp Gly Asp Val His Gly His Lys Phe Ser Val Arg Gly Glu
 20 25 30

Gly Glu Gly Asp Ala Asp Tyr Gly Lys Leu Glu Ile Lys Phe Ile Cys
 35 40 45

Thr Thr Gly Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr Phe
 50 55 60

Ser Tyr Gly Ile Gln Cys Phe Ala Arg Tyr Pro Glu His Met Lys Met
 65 70 75 80

Asn Asp Phe Phe Lys Ser Ala Met Pro Glu Gly Tyr Ile Gln Glu Arg
 85 90 95

Thr Ile Phe Phe Gln Asp Asp Gly Lys Tyr Lys Thr Arg Gly Glu Val
 100 105 110

Lys Phe Glu Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly Met
 115 120 125

Asp Phe Lys Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn
 130 135 140

Phe Asn Ser His Asn Val Tyr Ile Met Pro Asp Lys Ala Asn Asn Gly

145

150

155

160

Leu Lys Val Asn Phe Lys Ile Arg His Asn Ile Glu Gly Gly Gly Val
165 170 175

Gln Leu Ala Asp His Tyr Gln Thr Asn Val Pro Leu Gly Asp Gly Pro
180 185 190

Val Leu Ile Pro Ile Asn His Tyr Leu Ser Thr Gln Thr Ala Ile Ser
195 200 205

Lys Asp Arg Asn Glu Thr Arg Asp His Met Val Phe Leu Glu Phe Phe
210 215 220

Ser Ala Cys Gly His Thr His Gly Met Asp Glu Leu Tyr Lys
225 230 235

